

A new species (*Begonia giganticaulis*) of Begoniaceae from southern Xizang (Tibet) of China

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Abstract

Begonia giganticaulis, a huge new species in *Begonia* sect. *Platycentrum* of Begoniaceae from southern Xizang (Tibet) of China, is described. Morphologically, it is mostly similar to *B. longifolia* and *B. acetosella*, but clearly differs from the former mainly by its dioecious and taller plants, sparse hairs on abaxial veins, longer inflorescence, unique shape of fruits, and differs from the latter mainly by its late and longer flowering time, 6-tepals of female flower and 3-loculed ovary. The phylogenetic analyses also support the separation of the new species from other taxa. Based on the current data, its conservation status is assigned to Endangered (B2a) according to the IUCN Red List Categories and Criteria.

Keywords

Conservation status, molecular evidence, morphology, southern Tibet, taxonomy

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Introduction

Zangnan (southern Tibet) of China is located to the south of the Himalayas, including most parts of Cona, Lhünzê, Mêdog and Zayü counties, and some smaller parts of Nang and Mainling counties (Liu 2019). This region is very warm and rainy because of the southwest monsoon carrying heavy water and heat from the Indian Ocean. Owing to high average annual precipitation and high-proportion of forest coverage (Hao et al. 2010), the plant diversity is very high in Zangnan. However, this area still remains under-explored and needs more study in the future.

After a series of plant surveys recently, the authors have a better understanding of the diversity of *Begonia* in Tibet, particularly in its southern part (namely Zangnan) including Mêdog county. Up until now, 39 species and 4 varieties had been found in Tibet (Gu et al. 2007; Camfield and Hughes 2018; Tian et al. 2020) (Table 1). In addition, *Begonia limprichtii* Irmsch. (Irmscher 1922) was newly reported by Borah et al. (2021a) in southern Tibet, but this record is likely based on a wrong identification and further study is needed. Of these, 31 species and 3 varieties are distributed in Mêdog. Recently, after several field surveys in Mêdog, we found several new species and at least three natural hybrids. Here we described *Begonia giganticaulis* D.K.Tian & W.G.Wang sp. nov. from Mêdog, a new species of huge plant size, which is morphologically similar to both *B. longifolia* Blume (Blume 1827) and *B. acetosella* Craib (Craib 1912). The morphological differences of the three species are compared, and the new species is also supported by molecular evidence.

Material and methods

Morphological analysis

The field surveys were conducted on habitat, distribution, population size, morphology and specimen collection of the new species. Diagnosis of the morphological difference between the new species and its similar species was based on literature review, examination of herbarium specimens, and observation of both wild and cultivated plants.

Phylogenetic analysis

The treatment on sections of *Begonia* follows Shui et al. (2019). To ascertain the relationship of the new species within sect. *Platycentrum* (Klotzsch) A.DC. (de Candolle 1859), two female and three male individuals were sampled, and three individuals of *B. longifolia*, two individuals of *B. acetosella*, and three individuals of *B. acetosella* var. *hirtifolia* Irmsch. (Irmscher 1939) were sampled and sequenced. 13 taxa within sect. *Platycentrum* were selected based on Moonlight et al. (2018) to ascertain the phylogenetic relationship of the new species. *Begonia cavaleriei* H.Lév. (Léveillé 1909) from sect. *Coelocentrum* Irmsch. (Irmscher 1939) was used as outgroup. All

Table 1. A checklist of *Begonia* species in Tibet.

Species	Reference	County
<i>Begonia aborensis</i> Dunn	Dunn 1920	Médog
<i>Begonia acetosella</i> Craib	Craib 1912	Médog
<i>Begonia annulata</i> K.Koch	Koch 1857	Médog
<i>Begonia asperifolia</i> Irmsch.	Irmscher 1927	Bomê, Zayü, Lhünzê, Médog
<i>Begonia burkillii</i> Dunn	Dunn 1920	Médog
<i>Begonia cathcartii</i> Hook.f. & Thomson	Hooker 1855	Zayü
<i>Begonia dioica</i> Buch.-Ham. ex D.Don	Don 1825	Dinggyê
<i>Begonia difformis</i> (Irmsch.) W.C.Leong, C.I Peng & K.F.Chung	Leong et al. 2015	Médog
<i>Begonia flagellaris</i> Hara	Hara 1973	Gyirong, Nyalam
<i>Begonia flaviflora</i> var. <i>flaviflora</i> Hara	Hara 1970	Médog
<i>Begonia flaviflora</i> var. <i>gamblei</i> (Irmscher) Golding & Karegeannes	Golding and Karegeannes 1984	Médog
<i>Begonia gigantea</i> D.K.Tian & W.G. Wang sp. nov.	In this study	Médog
<i>Begonia grandis</i> Dryand.	Dryander 1791	Zayü
<i>Begonia griffithiana</i> (A.DC.) Warb.	Warburg 1894	Médog
<i>Begonia handelii</i> Irmsch.	Irmscher 1921	Médog
<i>Begonia hatacoa</i> Buch.-Ham. ex D.Don	Don 1825	Médog, Cona
<i>Begonia iridescens</i> Dunn	Dunn 1920	Médog, Zayü
<i>Begonia josephii</i> A.DC.	de Candolle 1859	Cona, Dinggyê, Lhün- zê, Médog, Yadong
<i>Begonia kekarmonyingensis</i> Taram, D.Borah & M.Hughes	Taram et al. 2021	Médog
<i>Begonia labordei</i> H.Lév.	Léveillé 1904	Zayü
<i>Begonia limprichtii</i> Irmsch. *(wrong identification, based on the distribu- tion and morphological characteristics)	Borah et al. 2021a	Médog
<i>Begonia longifolia</i> Blume	Blume 1827	Médog
<i>Begonia medogensis</i> J.W.Li, Y.H.Tan & X.H.Jin	Li et al. 2018	Médog
<i>Begonia megaloptera</i> A.DC.	de Candolle 1859	Zayü
<i>Begonia nepalensis</i> (A.DC.) Warb.	Warburg 1894	Cona
<i>Begonia ovatifolia</i> A.DC.	de Candolle 1859	Médog
<i>Begonia oyuniae</i> M.Taram & N.Krishna	Taram et al. 2020	Médog
<i>Begonia plamata</i> var. <i>plamata</i> D.Don	Don 1825	Médog
<i>Begonia palmata</i> var. <i>bowringiana</i> (Champion ex Benth) Golding & Karegeannes	Golding and Karegeannes 1984	Médog
<i>Begonia palmata</i> var. <i>hasiana</i> (Irmsch.) Golding & Kareg	Golding and Karegeannes 1984	Médog
<i>Begonia pasighatensis</i> D.Borah, Taram & Wahlsteen	Borah et al. 2021b	Médog
<i>Begonia picta</i> Sm.	Smith 1805	Gyirong, Médog, Nyalam
<i>Begonia pseudoheydei</i> Y.M.Shui & W.H.Chen	Chen et al. 2019	Médog
<i>Begonia rex</i> Putz.	Putzey 1857	Médog
<i>Begonia roxburghii</i> (Miq.) A.DC.	de Candolle 1864	Médog
<i>Begonia scintillans</i> Dunn	Dunn 1920	Médog
<i>Begonia shilendrae</i> Rekha Morris & P.D.McMillan	Morris and McMillan 2012	Cona
<i>Begonia sikkimensis</i> var. <i>sikkimensis</i> A.DC.	de Candolle 1859	Médog
<i>Begonia sikkimensis</i> var. <i>kamengensis</i> Rekha Morris, P.D.McMillan & Golding ex Golding	Golding 2009	Cona
<i>Begonia silletensis</i> Clarke	Clarke 1879	Médog
<i>Begonia tessaricarpa</i> C.B.Clarke	Clarke 1879	Médog
<i>Begonia thomsonii</i> A.DC.	de Candolle 1859	Médog
<i>Begonia xanthina</i> Hook.f.	Hooker 1852	Médog
<i>Begonia zhongyangiana</i> W.G.Wang et S.Z.Zhang	Wang et al. 2019	Médog

the voucher specimens were deposited in the herbarium of Xishuangbanna Tropical Botanical Garden, Chinese Academy of Sciences (HITBC). For DNA sequencing, the total genomic DNA was extracted from silica-dried leaves by a modified CTAB

Table 2. Sampled taxa and GenBank accession numbers of *Begonia giganticaulis* and the related taxa used for phylogenetic analysis.

Taxa	Collector, Voucher (Herbarium)	Origin	ITS	<i>rpL16</i>	<i>ndhA</i>	References
<i>Begonia acetosella</i> Craib	Wang, W.G., WWG004 (HITBC)	Mengla, Yunnan, China	MW690105	MW658199	MW658212	In this study
<i>B. acetosella</i> Craib	Wang, W.G., WWG005 (HITBC)	Mengla, Yunnan, China	MW690106	MW658200	MW658213	In this study
<i>B. acetosella</i> var. <i>hirtifolia</i> Irmsch.	Wang, W.G., WWG0261 (HITBC)	Ruili, Yunnan, China	MW690107	MW658201	—	In this study
<i>B. acetosella</i> var. <i>hirtifolia</i> Irmsch.	Wang, W.G., WWG0262 (HITBC)	Ruili, Yunnan, China	MW690108	MW658202	MW658214	In this study
<i>B. acetosella</i> var. <i>hirtifolia</i> Irmsch.	Wang, W.G., WWG0300 (HITBC)	Ruili, Yunnan, China	—	MW658203	MW658215	In this study
<i>B. aptera</i> Blume	—	—	AJ491196	—	JF756369	Chiang (unpublished); Thomas et al. (2011)
<i>B. balansana</i> Gagnep.	—	—	AF485091	KF707939	MH207051	Forrest and Hollingsworth (2003); Chung et al. (2014); Moonlight et al. (2018)
<i>B. cathayana</i> Hemsl.	—	—	AF280106	KF707948	KT599095	Yang et al. (unpublished); Chung et al. (2014); Zhao (unpublished)
<i>B. cavaleriei</i> H.Lév.	—	—	KF636430	KF707949	MK548079	Chung et al. (2014); Tong et al. (2019)
<i>B. decora</i> Stapf	—	—	KF636435	KF707956	JF756355	Chung et al. (2014); Thomas et al. (2011)
<i>B. giganticaulis</i> D.K.Tian & W.G.Wang	Wang, W.G., Li, Y.Y., Ma, X.D. & Shen, J.Y., WWG2014–1 (HITBC)	Médog, Tibet, China	MW690097	MW658191	MW658204	In this study
<i>B. giganticaulis</i> D.K.Tian & W.G.Wang	Wang, W.G., Li, Y.Y., Ma, X.D. & Shen, J.Y., WWG2015–1 (HITBC)	Médog, Tibet, China	MW690098	MW658192	MW658205	In this study
<i>B. giganticaulis</i> D.K.Tian & W.G.Wang	Wang, W.G., Li, Y.Y., Ma, X.D. & Shen, J.Y., WWG2015–2 (HITBC)	Médog, Tibet, China	MW690099	MW658193	MW658206	In this study
<i>B. giganticaulis</i> D.K.Tian & W.G.Wang	Wang, W.G., Li, Y.Y., Ma, X.D. & Shen, J.Y., WWG2014–3 (HITBC)	Médog, Tibet, China	MW690100	MW658194	MW658207	In this study
<i>B. giganticaulis</i> D.K.Tian & W.G.Wang	Wang, W.G., Li, Y.Y., Ma, X.D. & Shen, J.Y., WWG2014–2 (HITBC)	Médog, Tibet, China	MW690101	MW658195	MW658208	In this study
<i>B. handelii</i> Irmsch.	—	—	AF485093	KF707969	MH207176	Forrest and Hollingsworth (unpublished); Chung et al. (2014); Moonlight et al. (2018)
<i>B. hatacoa</i> Buch.-Ham. ex D.Don	—	—	AF485111	KF707970	JF756354	Forrest and Hollingsworth (unpublished); Chung et al. (2014); Thomas et al. (2011)
<i>B. longifolia</i> Blume	Wang, W.G., WWG001 (HITBC)	Mengla, Yunnan, China	MW690102	MW658196	MW658209	In this study
<i>B. longifolia</i> Blume	Wang, W.G., WWG002 (HITBC)	Mengla, Yunnan, China	MW690103	MW658197	MW658210	In this study

Taxa	Collector, Voucher (Herbarium)	Origin	ITS	<i>rpL16</i>	<i>ndhA</i>	References
<i>B. longifolia</i> Blume	Wang, W.G., WWG003 (HITBC)	Mengla, Yunnan, China	MW690104	MW658198	MW658211	In this study
<i>B. nepalensis</i> (A.DC.) Warb.	—	—	AY753726	—	MH207257	Tebbitt et al. (2006); Moonlight et al. (2018)
<i>B. obovoidea</i> Craib	—	—	—	—	JF756386	Thomas et al. (2011)
<i>B. pavonina</i> Ridl.	—	—	KF636472	KF708002	JF756356	Chung et al. (2014); Thomas et al. (2011)
<i>B. pedatifida</i> H.Lév.	—	—	MK541092	MK548068	MK548115	Tong et al. (2019)
<i>B. roxburghii</i> A.DC.	—	—	AF485092	—	JF756371	Forrest and Hollingsworth (2003); Thomas et al. (2011)
<i>B. versicolor</i> Irmsch.	—	—	AF485090	KF708023	JF756358	Forrest and Hollingsworth (unpublished); Chung et al. (2014); Thomas et al. (2011)

protocol (Doyle and Doyle 1987). The chloroplast DNA *rpL16* intron, *ndhA* intron and the nuclear ribosomal DNA internal transcribed spacer (nrITS) region were used to infer the phylogenetic relationship of the new species. The *rpL16* intron were amplified by the primer rpL16-F and rpL16-R and sequenced by the primer rpL16-R and Beg-rpL16 (Chung et al. 2014). For the amplification of the *ndhA* intron the primer *ndhAX1* and *ndhAX2* (Thomas et al. 2011) were used. The nrITS region was amplified and sequenced by the primer 51NT and 26S1Rev (Clement et al. 2004). The sampled sequences were downloaded from NCBI and accession numbers were listed in Table 2.

Sequences of each DNA region were aligned by MUSCLE online (<https://www.ebi.ac.uk/Tools/msa/muscle/>, Madeira et al. 2019) and adjusted manually when necessary. Indels were treated as gap. For testing the congruence within *rpL16* intron, *ndhA* intron and nrITS, the analysis of the incongruence length difference (ILD) was performed with 100 replicates under default heuristic search using PAUP v.4.0a (Swofford 2002) and the phylogenetic trees were constructed based on each dataset. The p value was 0.40 and no conflict among each phylogenetic trees, indicating the congruence among these datasets (Farris et al. 1994).

The parsimony analysis was conducted using PAUP v.4.0 b10 (Swofford 2002). The Maximum Parsimony (MP) analysis was run using a heuristic search with 1,000 replicates and tree-bisection-reconnection (TBR) with no reconnection limit. Bootstrap was used to assess the node support by 1000 replicates using TBR branch swapping. The Bayesian analysis was conducted using MrBayes v.3.1.2 (Ronquist and Huelsenbeck 2003) with 1,000,000 generations under the Markov chain Monte Carlo (MCMC) chains. The average standard deviation of split frequencies was 0.004210 after 1,000,000 generations. The consensus tree was constructed after burn-in 25% of the trees. The Posterior Probability (PP) was used to assess the branch supports.

Results

Taxonomic treatment

***Begonia giganticaulis* D.K.Tian & W.G.Wang, sp. nov.**

urn:lsid:ipni.org:names:77234844-1

Figs 1–4

巨型秋海棠

Type. CHINA. Xizang (Tibet) Autonomous Region: Mêdog county (墨脱县), Beibeng town (背崩乡), Baimu Xiri river (白母西日河), forest slope of river valley or water's edge along stream, 29°21'9"N, 95°11'21"E, elev. 1320 m, 10 September 2020, *Dai-Ke Tian, Fang Wen, Qing-Gong Mao, & Zhu Lu*, TDK4773-A (holotype CSH! Barcode number: 0180561, ♀)

Diagnosis. The new species is mostly similar to *B. longifolia* and *B. acetosella*, but clearly differs from the former mainly by its dioecious (vs. monoecious), taller (to 4 m vs. less than 2 m) plants, longer (vs. shorter) inflorescence, and unique shape of fruits, and differs from the latter mainly by its taller (to 4 m vs. less than 2 m) plants, late and longer (Jun. to Oct. vs. Mar. to Apr.) flowering time, longer (6–20 mm vs. 5–12 mm) pedicel, 6 (vs. 4) tepals of pistillate flower and 3 (vs. 4)-loculed ovary (Table 3, Fig. 3).

Description. **Herb** perennial, evergreen, to 4 m tall, dioecious. **Rhizome** short, stout, nearly unbranched, reddish brown, to 12 cm thick. **Stem** erect, reddish brown or green, glabrous, internodes to 5 cm thick, with many longitudinally fusiform whitish spots, cross section of stem often reddish brown, nodes notably enlarged, to 7 cm thick, with unequally oval to round whitish spots, many shrubby branches on the upper part of main stem. **Stipule** long-triangular, light green or pinkish green, 9–25 × 2–8 mm, glabrous, margin entire, dorsal ridge pinkish, apex acuminate with arista 4–6 mm long. **Petioles** green, pink to red, glabrous, 7–22 cm long, 1–3 mm thick. **Leaf blade** ovate-lanceolate to lanceolate, 4–19 × 0.8–8 cm, adaxial green, muriculate to nearly glabrous, adaxial veins slightly concave; abaxial greyish green, veins usually red, convex, main veins sparsely and obliquely strigose; base obliquely cordate, margin shallowly and remotely denticulate, apex long caudate; **Inflorescence** dichasial cyme, axillary, short, 3–5 cm long, unbranched to branched once, rachis glabrous, green, pinkish green to red, base usually red-brown, 7–15 mm long, 1–1.5 mm thick, 3–11 male flowers or 1–5 female per inflorescence. **Bract** often caducous, pinkish green, long triangular, glabrous, ca. 6 × 3 mm, apex acuminate; bracteoles smaller. **Staminate flower:** pedicel glabrous, white, whitish or pinkish green, 10–14 mm long, ca. 1 mm thick; corolla 18–24 mm in diameter; tepals 4, subequal, glabrous, outer 2, obovate, 9–14 × 6–9 mm, apex obtuse, adaxially white and middle-upper part abaxially pink, or pure white for some individuals, longitudinal veins unapparent; inner 2, pure white, obovate to obovate-lanceolate, 8–13 × 5–7 mm, apex obtuse; androecium nearly actinomorphic, ca. 5 mm long, 6–7 mm in diam; stamens 48–60, filaments free, 1–2 mm long; anther yellow, 2–3 mm long, apex obtuse or nearly so. **Pistillate flower:** pedicel white or green-white,



Figure 1. Habitat and large-sized plant of *Begonia giganticaulis* D.K.Tian & W.G.Wang, sp. nov. **A** habitat showing plants (arrows indicate) growing along stream bank **B** flowering plant growing along slope of valley **C** one of the tallest individuals with Dr. Dai-Ke Tian. (Photos **A** by Dai-Ke Tian **B** by Shi-Wei Guo **C** by Qing-Gong Mao).

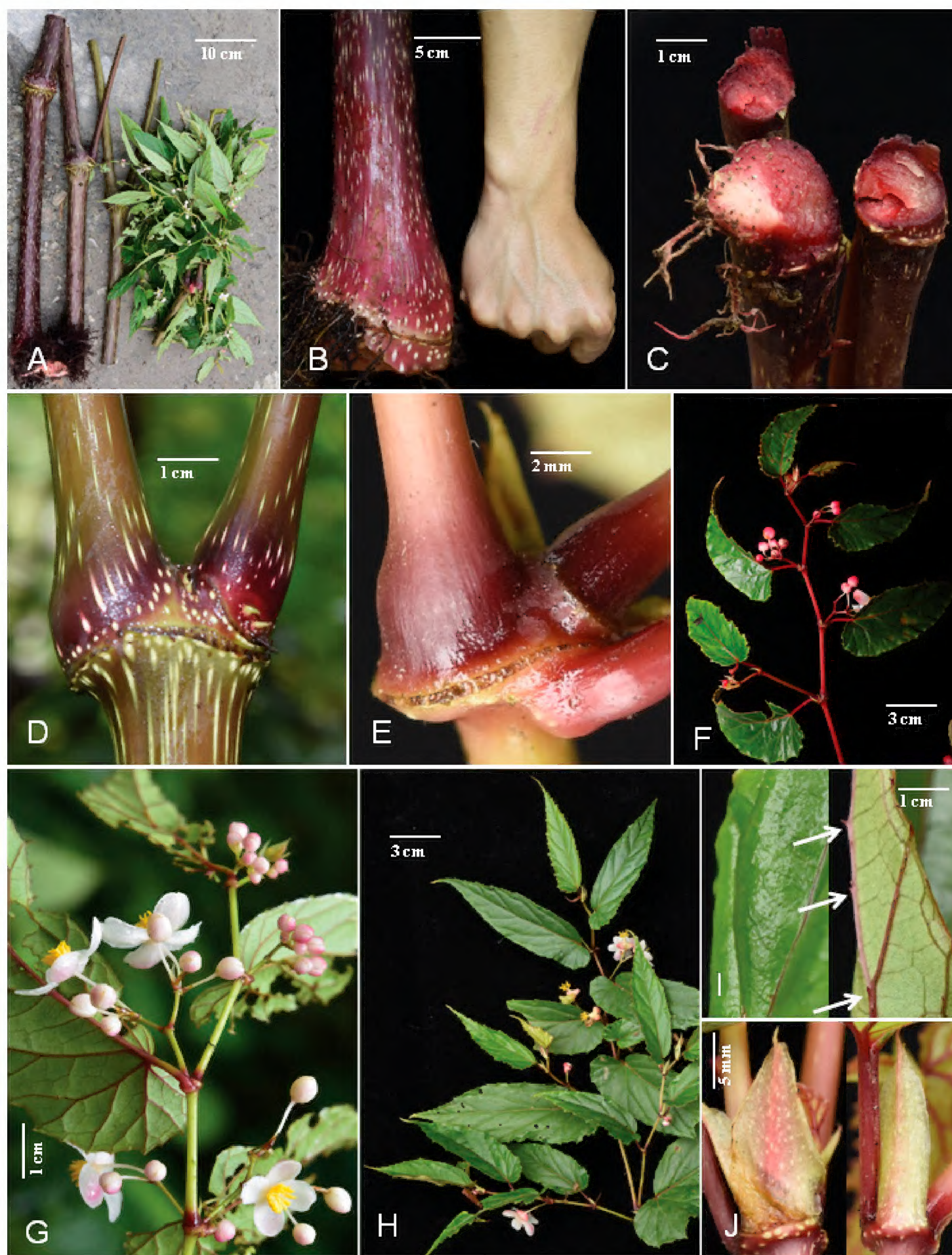


Figure 2. Morphology of *Begonia giganteaulis* D.K.Tian & W.G.Wang, sp. nov. **A** one of the single tallest plants cut into four sections **B** main stem base **C** stems showing colour of nodal cross-sections **D** main stem with much expanded node and whitish-green lines or spots **E** expanded node on terminal branch **F**, **G** male plant branches showing inflorescences and different colours **H** female branches **I** adaxially (left) nearly glabrous and abaxially sparse hairs on veins (right, arrows indicate) on blade surfaces **J** stipules showing shape and colour. (Photo **F** by Wen-Guang Wang; others by Dai-Ke Tian).

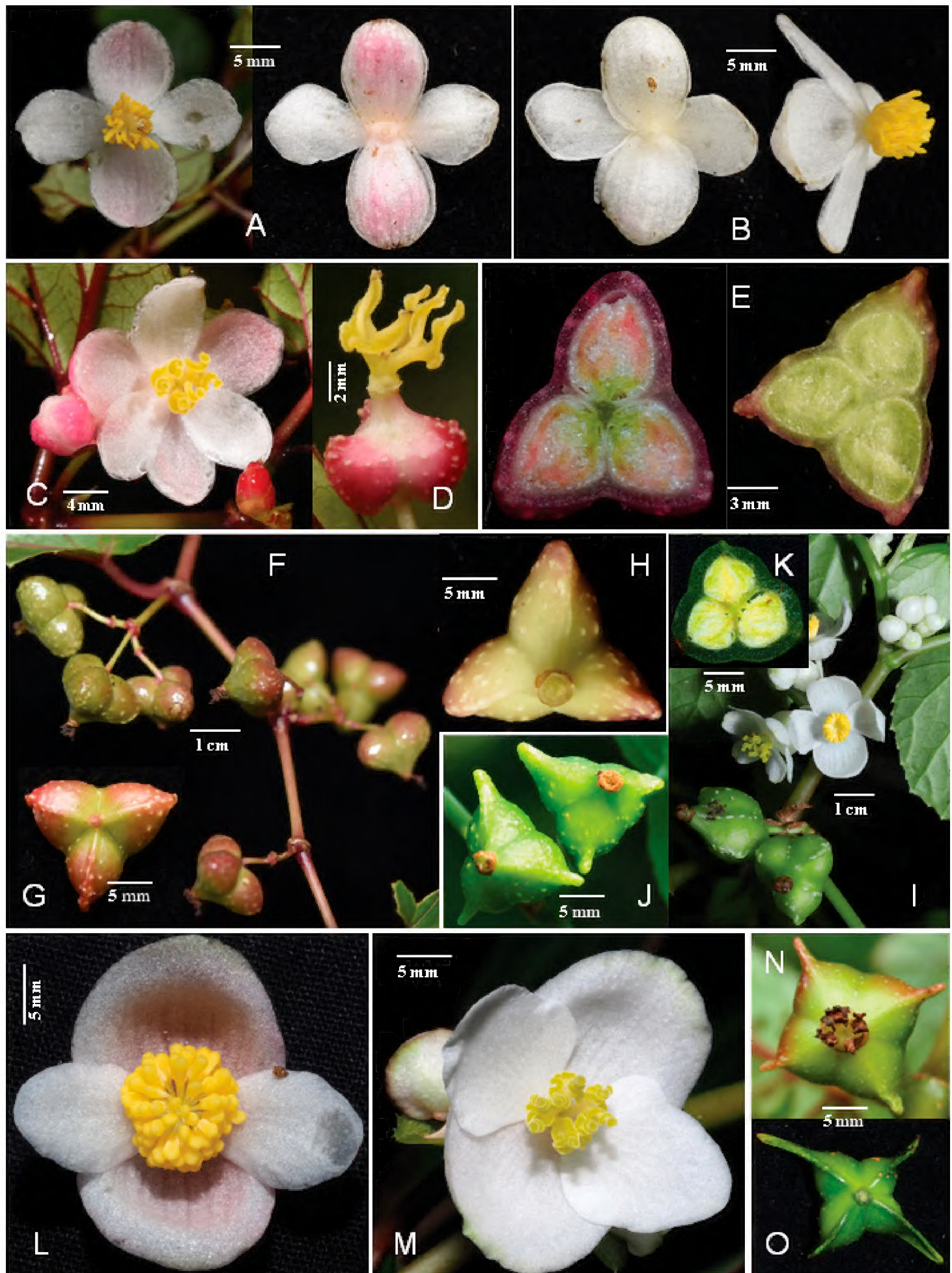


Figure 3. Flower and fruit morphology of *B. gigantea* compared with its close species *B. longifolia* and *B. acetosella* **A–H** *Begonia gigantea* **A** staminate flowers with pinkish outer tepals **B** staminate flowers with white tepals **C, D** pistillate flower **E** ovary sections showing different colour **F** fruits on branch **G, H** dorsal and front views of fruits **I–K** *B. longifolia* **I** flowering and fruiting branch **J** fruits showing short horns **K** ovary dissection **L–O** *B. acetosella* **L** staminate flower **M** Pistillate flower **N, O** fruits with short horns or wings. (Photos **C** by Shi-Wei Guo **E** (left) **L, M** & **O** by Wen-Guang Wang; others by Dai-Ke Tian).

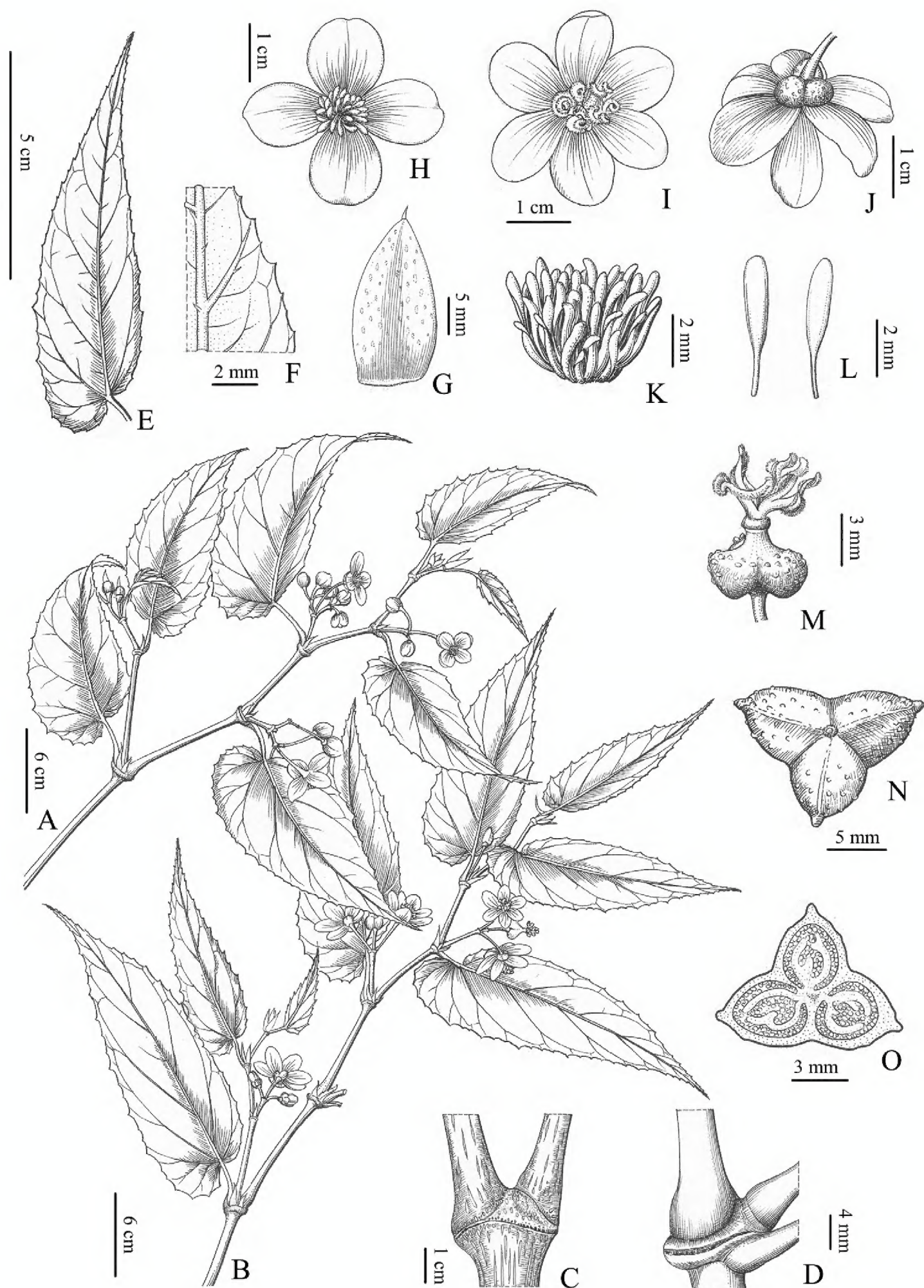


Figure 4. Illustration of *Begonia gigantea* D.K.Tian & W.G.Wang, sp. nov. (Drawn by Mr. Zhi-Min Li) **A** male flowering branches **B** female flowering branches **C** main stem line spots, much expanded node and internode base **D** expanded node and internode base on small upper branches **E** leaf blade **F** leaf (abaxial), showing sparse hairs on veins **G** stipule **H** staminate flowers **I, J** pistillate flower **K** side view of androecium **L** stamens **M** ovary and stigmas **N** fruit **O** dissection of ovary showing placentae.

Table 3. Morphological comparison of *Begonia giganticaulis*, *B. longifolia* and *B. acetosella*.

Character	<i>B. giganticaulis</i>	<i>B. longifolia</i>	<i>B. acetosella</i>
Plant			
sexuality	dioecious	monoecious	dioecious
height (m)	up to 4	less than 2	less than 2
Petiole length (cm)	0.7–7	1–12	1–10
Leaf blade surface	muriculate	glabrous to less muriculate	muriculate to hirsutulous
Inflorescence			
peduncle length (mm)	7–15	4–10	2–10
flower number	1–11	1–11(15)	1–3(5)
Tepal number of pistillate flower	6	6	4
Tepal colour	pinkish to white	white	pinkish to white
Ovary	3-loculed	3-loculed	4-loculed
Pedicel length (mm)			
male flower	10–20	5–12	8–12
female flower	6–12	5–12	5–10
Fruit horn or wing	none to rarely short crest	none to short crest or horns	short to long horns or wings
Flowering time	June–October	June–December	March–April

6–12 mm long, 0.8–1 mm thick; corolla 20–25 mm, tepals 6, rarely 4, glabrous, outer 3 (rarely 2), obovate or long obovate, thick and rigid, 12–18 × 7–10 mm, adaxial surface nearly white, distinctly concave, abaxially pink on middle-upper part, inner 3 (rarely 2), obovate-lanceolate to oblanceolate or long elliptical, slightly narrower than outer tepals, 10–19 × 6–8 mm, white, glabrous, apex obtuse; styles + stigmas 5 mm long, 7–8 mm wide; styles 3, free; stigmas yellow, nearly U-shaped, each side spirally twisted 1.5 circles; ovary pink or green, with white convex spots; placentation axile, 3-loculed, each placenta 2-branched. **Peduncle** green to pinkish green, glabrous, 8–12 mm long, ca. 1 mm thick. **Fruit** red, pink or green, glabrous, triangular-gyroscopic, 8–11 × 1–12 mm wide, concave between two placentas, wingless to occasionally short ridged, apex with beak 3–4 mm long. Flowering Jun.–Oct., fruiting Jul.–Dec.

Additional specimen examined. CHINA. **Xizang:** Mêdog County (墨脱县), Beibeng Town (背崩乡), Baimu Xiri River (白母西日河), forest slope of river valley or water's edge along stream, 29°21'9"N, 95°11'21"E, elev. 1320 m, 10 September 2020, *Dai-Ke Tian, Fang Wen, Qing-Gong Mao, & Zhu Lu* TDK4773-B (paratype CSH!, ♂); 29°20'0"N, 95°10'49"E, elev. 1110 m, 10 September 2020, *Dai-Ke Tian, Fang Wen, Qing-Gong Mao, & Zhu Lu* TDK4765-A, TDK4765-B, (paratype CSH!); 29°18'32"N, 95°10'38"E, elev. 980 m, 10 September 2020, *Dai-Ke Tian, Fang Wen, Qing-Gong Mao, & Zhu Lu* TDK4777 (paratype CSH!); near Ani Bridge (阿尼桥), 29°17'8.41"N, 95°10'3.23"E, elev. 810 m, 3 July 2020, *Wen-Guang Wang, You-Yun Li, Xing-Da Ma, & Jian-Yong Shen*, WWG 2014 (paratype, HITBC!), WWG 2015 (paratype HITBC!); elev. 1100 m, 16 September 1974, *anonymous* 2608 (paratype PE!); elev. 800–1400 m, 30 June 1980, *Wei-Lie Chen* 10809 (PE!); near No. 2 Bridge, 29°16'42"N, 95°10'49"E, elev. 810 m, 1 October 2017, *Dai-Ke Tian, Yan Xiao, Xin Zhong, Li-Zhi Tian & Zhu Lu* TDK3429 (paratype CSH!); Beibeng to Hanmi (汗密), elev. 840 m, 7 August 2010, South Tibet Expedition Team (藏南队), *Xiao-Hua Jin, Shu-Dong Zhang, Zhong-Yang Li, Bao-Cheng Wu, Xian-Yun Mu, Jing Li & Wei-Tao*

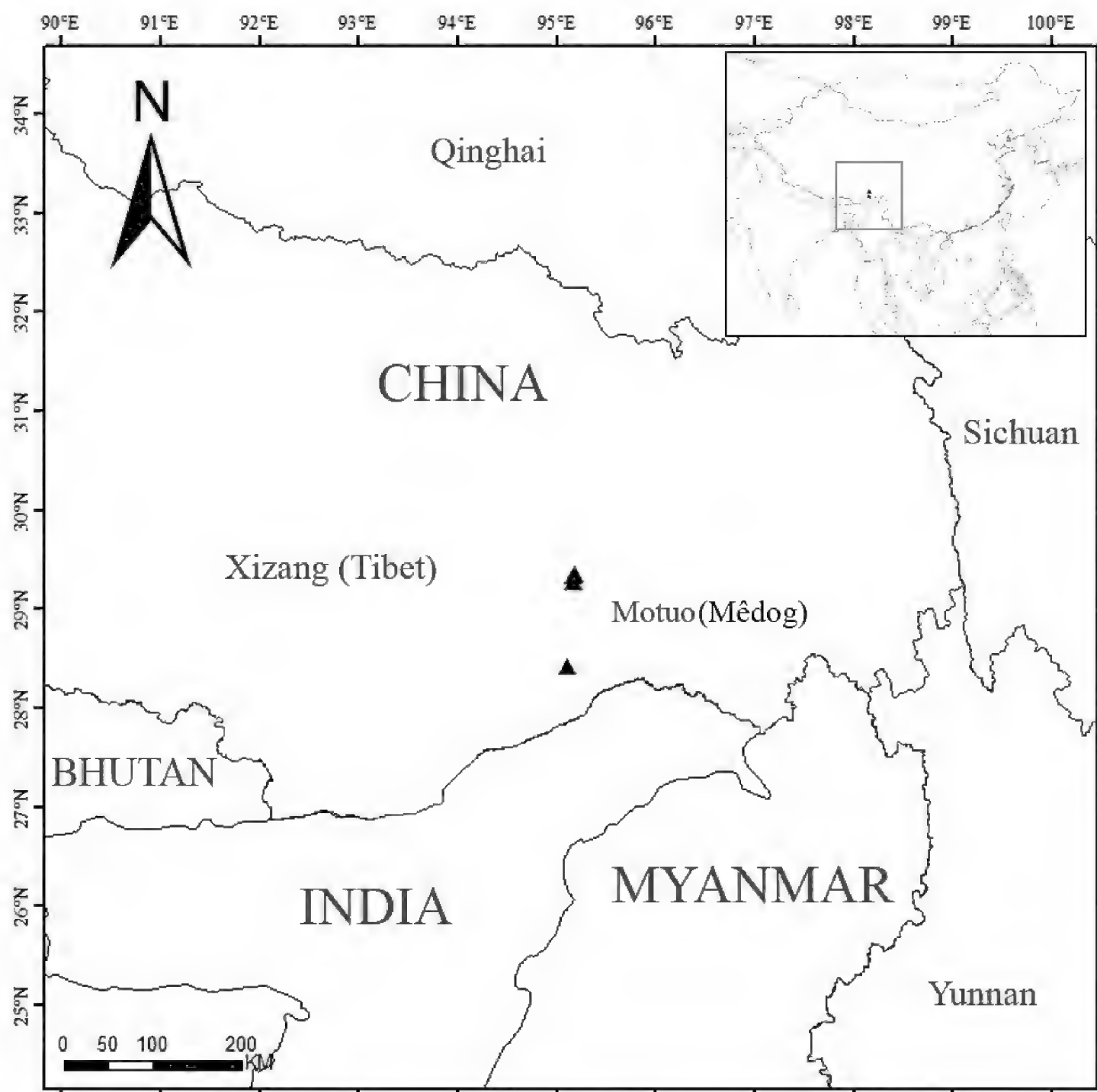


Figure 5. Distribution of *B. gigantecaulis* (triangles) from southern Xizang, China.

Jin, STET2304 (paratype PE!); Hanmi to Maniweng (马尼翁), elev. 800–1000 m, 6 August 1974, *Qingzang Team* 74-4114 (paratype PE!); elev. 1200 m, 24 June 1983, *Bo-Sheng Li & Shu-Zhi Chen* 05229 (paratype PE!); Maniweng to Ani Bridge, elev. 700–1000 m, 3 August 1972, *Tibet Expedition Team, Institute of Biology* 1631 (paratype HNWP!).

Distribution and habitat. Currently known from at least two localities in Mêdog, southern Xizang (Tibet), China (Fig. 5). It grows on the slopes under forest along streams, elevation 450–1400 m.

Conservation status. *Begonia gigantecaulis* is currently found in at least two localities in Mêdog of Tibet. Additional populations might be discovered when more surveys are conducted in China-India border region. However, based on current data, it should be categorised as Endangered: B2a (IUCN 2019) due to < 500 km² area of occupancy with severely fragmented habitat consisting of < 5 known populations totally under 1000 individuals by estimation.

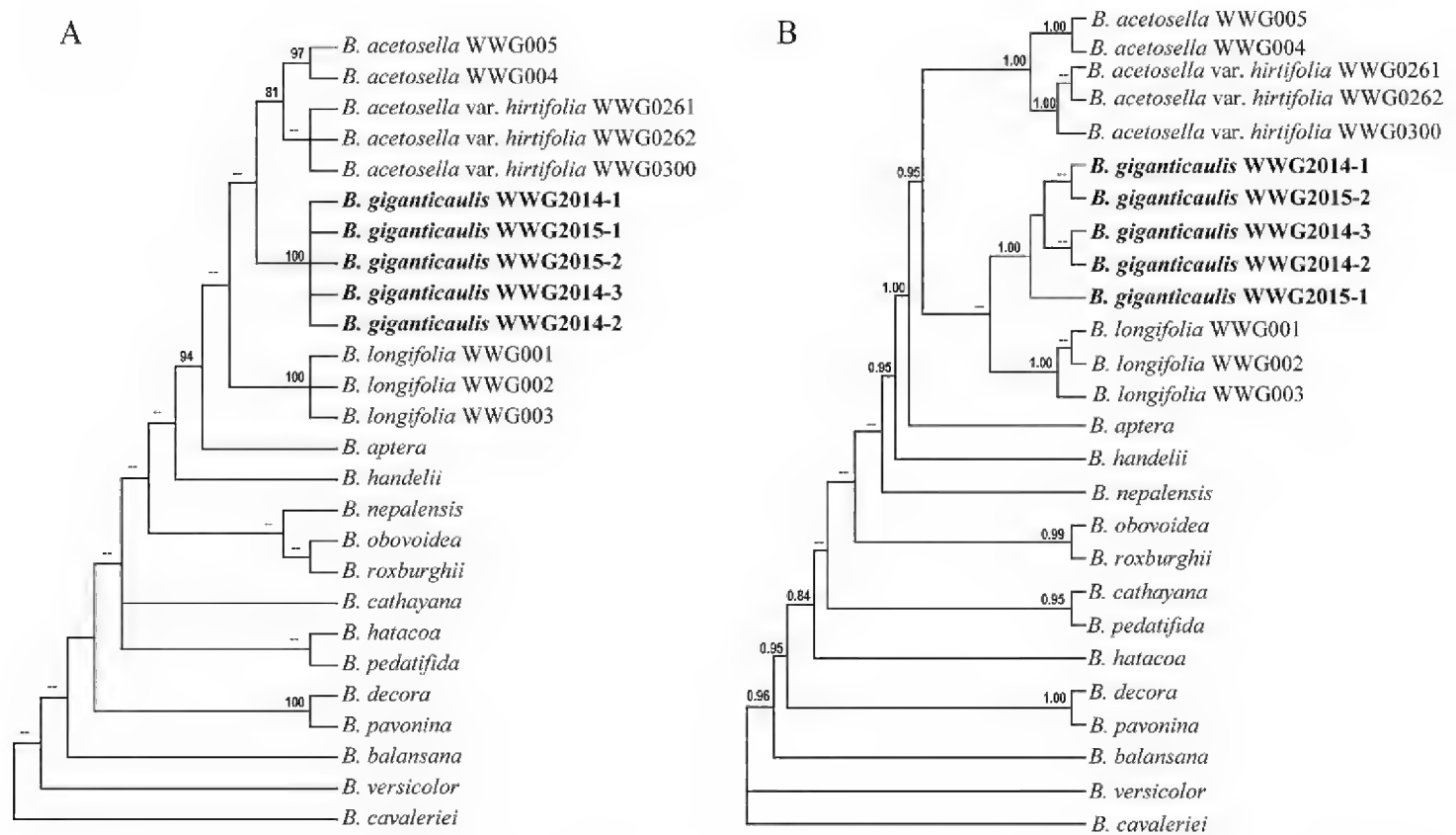


Figure 6. Phylogenetic tree inferred by MP **A** and BI **B** analyses based on the combined matrix of two plastid loci (*rpL16* intron and *ndhA* intron) and nuclear ITS. Maximum parsimony bootstrap **A** and Bayesian inference posterior probability values **B** are labelled on the branches; when the number is below 80 and 0.80 in maximum parsimony bootstrap and Bayesian inference posterior probability, respectively, the branches are labelled—.

Etymology. The specific epithet refers to the huge (very tall and thick stem) plant size of the new species, which is the tallest begonia in Asia.

Molecular systematic relationship

We obtained 12 nrITS, 13 *rpL16* intron, and 13 *ndhA* intron of the new species and related *Begonia* taxa. In order to reconstruct the phylogenetic relationship of the new species, 13 taxa within sect. *Platycentrum* were included and *B. cavaleriei* from sect. *Coelocentrum* was selected as outgroup. In total, the matrix was composed of 26 accessions and contained the 962 bp *rpL16* intron, the 1109 bp *ndhA* intron and the 672 bp nrITS sequence. Of the total 2743 characters, 132 were parsimony informative.

Based on MP analysis, the new species was clustered with *B. acetosella* and *B. acetosella* var. *hirtifolia* (Fig. 6A), while it was clustered with *B. longifolia* under BI analysis (Fig. 6B). Both MP and BI analyses showed that all five individuals of the new species were clustered together and separated from other taxa (Fig. 6A, BS: 100%; Fig. 6B, PP:1.00).

Notes. – The earliest specimen of *Begonia giganticaulis* was collected in 1972 between Maliweng and Ani Bridge, Mêdog, Tibet, China. This species is similar to *B. acetosella* in appearance when its flowers are unavailable for observation, therefore, it was misidentified (24 June 1983, *Bo-Sheng Li & Shu-Zhi Chen 05229*, PE! was wrongly identified as *B. acetosella* by C.Z. Gu in March 2004). Also, due to its high similarity to *B. longifolia* particularly in morphology of flowers and fruits, *B. giganticaulis* was

wrongly labelled as *B. longifolia* by Morris (2010) who found this species in southern Mêdog county.

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